



```
... -----} p35-(23)-p10 4173266 Gap 1.4 bits  
... -----| p35-p10 4173266 total 4.9 bits
```

Sequence alignment of nine proteins. The top sequence is the reference protein (4173420) and the bottom sequence is the query protein (4173410). Red dots indicate identical amino acids. Two purple boxes highlight regions of interest: one from position 100 to 120 and another from position 200 to 220. Below the alignment, two p-values are shown: 6.2 bits for the first box and 6.6 bits for the second.

A diagram of a DNA sequence. The top strand is labeled 5' \* g t a g g t 3'. The bottom strand is labeled 5' - - - - - 3'. A red box highlights the 6th nucleotide position, where the top strand has a 'g' and the bottom strand has a 't'.